



# SEQUENCE LISTING

<110> BROWNING, JEFFREY  
AMBROSE, CHRISTINE  
MACKAY, FABIENNE  
TSCHOPP, JURG  
SCHNEIDER, PASCAL

<120> BAFF, INHIBITORS THEREOF AND THEIR USE IN THE  
MODULATION OF B-CELL RESPONSE

<130> 08201.0024-00000

<140> 09/911,777

<141> 2001-07-24

<150> 60/143,228

<151> 2001-07-09

<150> PCT/US00/01788

<151> 2000-01-25

<150> 60/117,169

<151> 1999-01-25

<160> 26

<170> PatentIn Ver. 2.1

<210> 1

<211> 285

<212> PRT

<213> Homo sapiens

<400> 1

Met Asp Asp Ser Thr Glu Arg Glu Gln Ser Arg Leu Thr Ser Cys Leu  
1 5 10 15

Lys Lys Arg Glu Glu Met Lys Leu Lys Glu Cys Val Ser Ile Leu Pro  
20 25 30

Arg Lys Glu Ser Pro Ser Val Arg Ser Ser Lys Asp Gly Lys Leu Leu  
35 40 45

Ala Ala Thr Leu Leu Leu Ala Leu Leu Ser Cys Cys Leu Thr Val Val  
50 55 60

Ser Phe Tyr Gln Val Ala Ala Leu Gln Gly Asp Leu Ala Ser Leu Arg  
65 70 75 80

Ala Glu Leu Gln Gly His His Ala Glu Lys Leu Pro Ala Gly Ala Gly  
85 90 95

Ala Pro Lys Ala Gly Leu Glu Glu Ala Pro Ala Val Thr Ala Gly Leu  
100 105 110

Lys Ile Phe Glu Pro Pro Ala Pro Gly Glu Gly Asn Ser Ser Gln Asn  
115 120 125

Ser Arg Asn Lys Arg Ala Val Gln Gly Pro Glu Glu Thr Val Thr Gln  
130 135 140

Asp Cys Leu Gln Leu Ile Ala Asp Ser Glu Thr Pro Thr Ile Gln Lys  
145 150 155 160

Gly Ser Tyr Thr Phe Val Pro Trp Leu Leu Ser Phe Lys Arg Gly Ser  
165 170 175

Ala Leu Glu Glu Lys Glu Asn Lys Ile Leu Val Lys Glu Thr Gly Tyr  
180 185 190

Phe Phe Ile Tyr Gly Gln Val Leu Tyr Thr Asp Lys Thr Tyr Ala Met  
195 200 205

Gly His Leu Ile Gln Arg Lys Lys Val His Val Phe Gly Asp Glu Leu  
210 215 220

Ser Leu Val Thr Leu Phe Arg Cys Ile Gln Asn Met Pro Glu Thr Leu  
225 230 235 240

Pro Asn Asn Ser Cys Tyr Ser Ala Gly Ile Ala Lys Leu Glu Glu Gly  
245 250 255

Asp Glu Leu Gln Leu Ala Ile Pro Arg Glu Asn Ala Gln Ile Ser Leu  
260 265 270

Asp Gly Asp Val Thr Phe Phe Gly Ala Leu Lys Leu Leu  
275 280 285

<210> 2

<211> 309

<212> PRT

<213> Mus sp.

<400> 2

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Gln	Lys	Glu	Glu	Gly	Ala	Trp	Phe	Gly	Ile	Cys	Arg	Asp	Gly	Arg	Leu		
		35					40					45					
Leu	Ala	Ala	Thr	Leu	Leu	Leu	Ala	Leu	Leu	Ser	Ser	Ser	Phe	Thr	Ala		
	50					55					60						
Met	Ser	Leu	Tyr	Gln	Leu	Ala	Ala	Leu	Gln	Ala	Asp	Leu	Met	Asn	Leu		
65				70					75					80			
Arg	Met	Glu	Leu	Gln	Ser	Tyr	Arg	Gly	Ser	Ala	Thr	Pro	Ala	Ala	Ala		
			85					90						95			
Gly	Ala	Pro	Glu	Leu	Thr	Ala	Gly	Val	Lys	Leu	Leu	Thr	Pro	Ala	Ala		
		100						105					110				
Pro	Arg	Pro	His	Asn	Ser	Ser	Arg	Gly	His	Arg	Asn	Arg	Arg	Ala	Phe		
		115					120					125					
Gln	Gly	Pro	Glu	Glu	Thr	Glu	Gln	Asp	Val	Asp	Leu	Ser	Ala	Pro	Pro		
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Ala	Pro	Cys	Leu	Pro	Gly	Cys	Arg	His	Ser	Gln	His	Asp	Asp	Asn	Gly		
145					150					155				160			
Met	Asn	Leu	Arg	Asn	Ile	Ile	Gln	Asp	Cys	Leu	Gln	Leu	Ile	Ala	Asp		
				165					170					175			
Ser	Asp	Thr	Pro	Thr	Ile	Arg	Lys	Gly	Thr	Tyr	Thr	Phe	Val	Pro	Trp		
			180					185					190				
Leu	Leu	Ser	Phe	Lys	Arg	Gly	Asn	Ala	Leu	Glu	Glu	Lys	Glu	Asn	Lys		
		195					200					205					
Ile	Val	Val	Arg	Gln	Thr	Gly	Tyr	Phe	Phe	Ile	Tyr	Ser	Gln	Val	Leu		
	210					215					220						

Tyr Thr Asp Pro Ile Phe Ala Met Gly His Val Ile Gln Arg Lys Lys  
 225 230 235 240

Val His Val Phe Gly Asp Glu Leu Ser Leu Val Thr Leu Phe Arg Cys  
 245 250 255

Ile Gln Asn Met Pro Lys Thr Leu Pro Asn Asn Ser Cys Tyr Ser Ala  
 260 265 270

Gly Ile Ala Arg Leu Glu Glu Gly Asp Glu Ile Gln Leu Ala Ile Pro  
 275 280 285

Arg Glu Asn Ala Gln Ile Ser Arg Asn Gly Asp Asp Thr Phe Phe Gly  
 290 295 300

Ala Leu Lys Leu Leu  
 305

<210> 3  
 <211> 102  
 <212> PRT  
 <213> Homo sapiens

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 Val Thr Gln Asp Cys Leu Gln Leu Ile Ala Asp Ser Glu Thr Pro Thr  
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Ile Gln Lys Gly Ser Tyr Thr Phe Val Pro Trp Leu Leu Ser Phe Lys  
 20 25 30

Arg Gly Ser Ala Leu Glu Glu Lys Tyr Gly Gln Val Leu Tyr Thr Asp  
 35 40 45

Lys Thr Tyr Ala Met Gly His Leu Ile Gln Arg Lys Lys Val His Val  
 50 55 60

Phe Gly Asp Glu Leu Ser Asn Asn Ser Cys Tyr Ser Ala Gly Ile Ala  
 65 70 75 80

Lys Leu Glu Glu Gly Asp Glu Leu Gln Leu Ala Ile Pro Arg Glu Asn  
 85 90 95

Ala Gln Ile Ser Leu Asp  
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<210> 4  
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 <212> PRT  
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<400> 4

Lys Gln His Ser Val Leu His Leu Val Pro Ile Asn Ala Thr Ser Lys  
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 Asp Asp Ser Asp Val Thr Glu Val Met Trp Gln Pro Ala Leu Arg Arg  
 20 25 30  
 Gly Arg Gly Leu Gln Ala Gln Tyr Ser Gln Val Leu Phe Gln Asp Val  
 35 40 45  
 Thr Phe Thr Met Gly Gln Val Val Ser Arg Glu Gly Gln Gly Arg Ala  
 50 55 60  
 Tyr Asn Ser Cys Tyr Ser Ala Gly Val Phe His Leu His Gln Gly Asp  
 65 70 75 80  
 Ile Leu Ser Val Ile Ile Pro Arg Ala Arg Ala Lys Leu Asn Leu Ser  
 85 90 95

<210> 5  
 <211> 104  
 <212> PRT  
 <213> Homo sapiens

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 Gln Leu Gln Trp Leu Asn Arg Arg Ala Asn Ala Leu Leu Ala Asn Gly  
 20 25 30  
 Val Tyr Ser Gln Val Leu Phe Lys Gly Gln Gly Cys Pro Ser Thr His  
 35 40 45  
 Val Leu Leu Thr His Thr Ile Ser Arg Ile Ala Val Ser Tyr Gln Thr  
 50 55 60  
 Glu Gly Ala Glu Ala Lys Pro Trp Tyr Glu Pro Ile Tyr Leu Gly Gly  
 65 70 75 80  
 Val Phe Gln Leu Glu Lys Gly Asp Arg Leu Ser Ala Glu Ile Asn Arg  
 85 90 95  
 Pro Asp Tyr Leu Asp Phe Ala Glu  
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<210> 6  
 <211> 97  
 <212> PRT  
 <213> Homo sapiens

<400> 6  
 Glu Leu Arg Lys Val Ala His Leu Thr Gly Lys Ser Asn Ser Arg Ser  
 1 5 10 15

Met Pro Leu Glu Trp Glu Asp Thr Tyr Gly Ile Val Leu Leu Ser Gly  
20 25 30

Val Lys Tyr Ser Lys Val Tyr Phe Arg Gly Gln Ser Cys Asn Asn Leu  
35 40 45

Pro Leu Ser His Lys Val Tyr Met Arg Asn Ser Lys Tyr Pro Gln Met  
50 55 60

Trp Ala Arg Ser Ser Tyr Leu Gly Ala Val Phe Asn Leu Thr Ser Ala  
65 70 75 80

Asp His Leu Tyr Val Asn Val Ser Glu Leu Ser Leu Val Asn Phe Glu  
85 90 95

Glu

<210> 7  
<211> 102  
<212> PRT  
<213> Homo sapiens

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Thr Leu Lys Pro Ala Ala His Leu Ile Gly Asp Pro Ser Lys Gln Asn  
1 5 10 15

Ser Leu Leu Trp Arg Ala Asn Thr Asp Arg Ala Phe Leu Gln Asp Gly  
20 25 30

Phe Tyr Ser Gln Val Val Phe Ser Gly Lys Ala Tyr Ser Pro Lys Ala  
35 40 45

Thr Ser Ser Pro Leu Tyr Leu Ala His Glu Val Gln Leu Phe Ser Ser  
50 55 60

Gln Tyr Pro Phe Pro Trp Leu His Ser Met Tyr His Gly Ala Ala Phe  
65 70 75 80

Gln Leu Thr Gln Gly Asp Gln Leu Ser Thr His Thr Asp Gly Ile Pro  
85 90 95

His Leu Val Leu Ser Phe  
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<210> 8  
<211> 109  
<212> PRT  
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Glu Ala Gln Pro Phe Ala His Leu Thr Ile Asn Ala Thr Asp Ile Pro  
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Ser Gly Ser His Lys Val Ser Leu Ser Ser Trp Tyr His Asp Arg Gly  
20 25 30

Trp Gly Lys Ile Ser Asn Met Tyr Ala Asn Ile Cys Phe Arg His His  
           35                                  40                                  45  
 Glu Thr Ser Gly Asp Leu Ala Thr Glu Tyr Leu Gln Leu Met Val Tyr  
           50                                  55                                  60  
 Val Thr Lys Thr Ser Ile Lys Ile Pro Ser Glu Phe His Phe Tyr Ser  
           65                                  70                                  75                                  80  
 Ile Asn Val Gly Gly Phe Phe Lys Leu Arg Ser Gly Glu Glu Ile Ser  
                                   85                                  90                                  95  
 Ile Glu Val Ser Asn Pro Ser Leu Leu Asp Pro Asp Gln  
                                   100                                  105

<210> 9  
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<400> 9  
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<210> 10  
 <211> 30  
 <212> DNA  
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<400> 10  
 gacaagcttg ccaccatgga tgactccaca 30

<210> 11  
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<400> 11  
 actagtcaca gcagtttcaa tgc 23

<210> 12  
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<400> 12  
 ctgcagggtc cagaagaaac ag 22

<210> 13  
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<400> 20  
taagaatgcg gccgcgggat cacgcactcc agcaa 35

<210> 21  
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<400> 21  
gcagtttcac agcgatgtcc t 21

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<400> 22  
gtctccgttg cgtgaaatct g 21

<210> 23  
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<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Illustrative motif

<400> 23  
Arg Asn Lys Arg  
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<210> 24  
<211> 4  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Illustrative motif

<400> 24  
Arg Lys Arg Arg  
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<210> 25  
<211> 4  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Illustrative  
motif

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Arg Pro Arg Arg  
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<210> 26  
<211> 4  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Illustrative  
motif

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<222> (2)  
<223> Any amino acid

<220>  
<221> MOD\_RES  
<222> (3)  
<223> Lys or Arg

<400> 26  
Arg Xaa Xaa Arg  
1